## SEQUENCE LISTING





<110> Innogenetics N.V.

<120> New sequences of hepatitis C virus genotypes for diagnosis, prophylaxis and therapy

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- <212> DNA
- <213> hepatitis C virus
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Ser Phe Gly Asn Thr Ile Thr Cys Tyr Ile Lys Ala Thr Ala Ala Ala		
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75

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180

240

300

360

420

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540

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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45 `

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Thr Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Ala Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val

Gly Ala Thr Thr Ala Ser Ile Arg Arg His Val Asp Ile Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

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Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly 165 170 175

His Arg Met Ala 180

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<212> PRT

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Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly 20 25 30

Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Thr Pro Gly Cys Ile Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Lys Tyr Val

100 105 110

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly 115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160

Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly 165 170 175

His Arg Met Ala 180

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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Thr Pro Gly Cys Ile Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Lys Tyr Val

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly 115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160

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His Arg Met Ala 180

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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85. 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val
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Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 135 130 140 Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 150 145 Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly

His Arg Met Ala 180

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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val

Ile Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val 100 105 110

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly 115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

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cggc	cta	ctg d	ttgt	cagt	c gg	gctgc	gtts	, taa	atcgt	ggg	tcat	catc	gag	ctggg	ggggca
agco	ggc	act o	gtto	caga	ac aa	aggag	gtgt	tgt:	atca	aca	gtad	gate	gag	atgga	aggagt
gctc	gca	agc o	gccc	cata	ac at	cgaa	caaç	g cto	aggt	aat	agco	ccaco	cag	ttcaa	aggaga
aaat	cct	tgg a	actgo	etgca	ag cg	gagco	cacco	aac	caaca	agc	tgt	catte	gag	cccgt	taatag
cttc	caa	ctg g	gcaaa	agct	t ga	aaaco	ttct	ggd	cacaa	agca	t				
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<400	>	32													
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Cys	Met	Ser	Ala 20	Asp	Leu	Glu	Val	Thr 25	Thr	Ser	Thr	Trp	Val 30	Leu	Leu
Gly	Gly	Val 35	Leu	Ala	Ala	Leu	Ala 40	Ala	Tyr	Cys	Leu	Ser 45	Val	Gly	Cys
Val	Val 50	Ile	Val	Gly	His	Ile 55	Glu	Leu	Gly	Gly	Lys 60	Pro	Ala	Leu	Val
Pro 65	Asp	Lys	Glu	Val	Leu 70	Tyr	Gln	Gln	Tyr	Asp 75	Glu	Met	Glu	Glu	Cys 80
Ser	Gln	Ala	Ala	Pro 85	Tyr	Ile	Glu	Gln	Ala 90	Gln	Val	Ile	Ala	His 95	Gln
Phe	Lys	Glu	Lys 100	Ile	Leu	Gly	Leu	Leu 105	Gln	Arg	Ala	Thr	Gln 110	Gln	Gln

Ala Val Ile Glu Pro Val Ile Ala Ser Asn Trp Gln Lys Leu Glu Thr

120

125

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tgttgcttgg aggggtcctc gcggccctag

60

120

115

Phe Trp His Lys His

<210><211><212><212><213>	33 401 DNA	atitis C vi	rue
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gctcgcaagc cgccccatac atcgaacaag ctcaggtaat agcccaccag ttcaaggaga 300
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cttccaactg gcaaaagctt gaaaccttct ggcacaagca t 401

<210> 34 <211> 133 <212> PRT

<213> hepatitis C virus

<400> 34

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Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu 20 25 30

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35 40 45

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Leu Val 50 55 60

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys 70 75 80

Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln 85 90 95

Phe Lys Glu Lys Ile Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln Gln 100

Ala Val Ile Glu Pro Val Ile Ala Ser Asn Trp Gln Lys Leu Glu Thr 115

Phe Trp His Lys His

Phe Trp His Lys His 130

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<213> hepatitis C virus

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cggcctactg cttgtcagtc ggttgtgttg tgattgtggg tcatatcgag ctggggggca 180
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gctcacaagc tgccccatat atcgaacaag ctcaggtaat agctcaccag ttcaagggaa 300
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<210> 36 <211> 133 <212> PRT <213> hepatitis C virus

<400> 36

Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala 1 5 10 15

Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu 20 25 30

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35 40 45

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val 50 55 60

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Gly Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln 105 100 Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 120 115 Phe Trp His Lys His 130 <210> 37 <211> 401 DNA <212> <213> hepatitis C virus <400> 37 tecaaaatga aatetgettg acacacecca teacaaaata cateatggea tgeatgteag 60 ctgatctgga agtaaccacc agcacctggg ttttgcttgg aggggtcctc gcggccctag 120 eggectactg ettgteagte ggttgtgttg tgattgtggg teatategag etggggggea 180 agccggcaat cgttccagac aaagaggtgt tgtatcaaca atacgatgag atggaagagt 240 gctcacaagc tgccccatat atcgaacaag ctcaggtgat agctcaccag ttcaaggaaa 300 aagtoottgg attgotgcag cgagccaccc aacaacaagc tgtcattgag cccatagtaa 360 401 ctaccaactg gcaaaagctt gaggcctttt ggcacaagca t <210> 38 <211> 133 PRT <212> <213> hepatitis C virus <400> .38 Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala

Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu 25

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val 50

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys 70

Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln 85 90 95

Phe Lys Glu Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln Gln 100 105 110

Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 115 120 125

Phe Trp His Lys His 130

<210> 39 <211> 401 <212> DNA

<213> hepatitis C virus

<400> 39
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cggcctactg cttgtcagtc ggttgtgttg tgattgtggg tcatatcgag ctggggggca 180
agccggcaat cgttccagac aaagaggtgt tgtatcaaca atacgatgag atggaagagt 240
gctcacaagc tgccccatat atcgaacaag ctcaggtaat agctcaccag ttcaaggaaa 300
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<210> 40 <211> 133 <212> PRT <213> hepatitis C virus

<400> 40

Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala

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Cys Met Ser Ala	Asp Leu Glu V	Val Thr Th 25	nr Ser Thr	Trp Val Le	u Léu
Gly Gly Val Leu 35		Ala Ala Ty 40	r Cys Leu	Ser Val Gl 45	y Cys
Val Val Ile Val	Gly His Ile (	Glu Leu Gl	ly Gly Lys 60	Pro Ala Il	e Val
Pro Asp Lys Glu	Val Leu Tyr ( 70	Gln Gln Ty	yr Asp Glu 75	Met Glu Gl	u Cys 80
Ser Gln Ala Ala	Pro Tyr Ile (	Glu Gln Al 90	la Gln Val O	Ile Ala Hi	s Gln
Phe Lys Glu Lys		Leu Leu G 105	ln Arg Ala	Thr Gln Gl	in Gln
Ala Val Ile Glu 115		Thr Thr A	sn Trp Gln	Lys Leu Gl	lu Ala
Phe Trp His Lys	s His				
<210> 41 <211> 509 <212> DNA <213> hepatit:	is C virus				
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aggacgtcaa gtto					
ggggccctag gat					
gacggcgtca gcc	tattccc aaggcg	gegee agee	cacggg ccg	gtcctgg gg	
ggtacccttg gcc	cctttac gccaat	gagg gcct	cgggtg ggc	agggtgg ct	gctctccc 300

ctcgaggetc teggeetaat tggggeecca atgaeeceeg gegaaaateg egtaatttgg

gtaaggtcat cgatacccta acgtgcggat tcgccgatct catggggtat atcccgctcg

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360

420

<210> 42

<211> 169

<212> PRT

<213> hepatitis C virus

<400> 42

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro 100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu 165

<210> 43

<211> 509

<212> DNA

## <213> hepatitis C virus

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ggggccctag	gatgggtgtg	cgcgcgactc	ggaagacttc	ggaacggtcg	caaccccgtg	180
gacggcgtca	gcctattccc	aaggcgcgcc	agcccacggg	ccggtcctgg	ggtcaacccg	240
ggtacccttg	gcccctttac	gccaatgagg	gcctcgggtg	ggcagggtgg	ctgctctccc	300
ctcgaggctc	tcggcctaat	tggggcccca	atgacccccg	gcgaaaatcg	cgtaatttgg	360
gtaaggtcat	cgatacccta	acgtgcggat	tcgccgatct	catggggtat	atcccgctcg	420
taggcggccc	cattgggggc	gtcgcaaggg	ctctcgcaca	cggtgtgagg	gtccttgagg	480
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<210> 44

<211> 169

<212> PRT

<213> hepatitis C virus

<400> 44

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp

85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu 165

<210> 45

<211> 580

<212> DNA

<213> hepatitis C virus

<400> 45

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<210> 46

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 46

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Ile Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

للم المرابع

Glu	Asp	Gly 35	Val	Asn	Tyr	Ala	Thr 40	Gly	Asn	Leu	Pro	Gly 45	Cys	Ser	Phe			
Ser	Ile 50	Phe	Ile	Leu	Ala	Leu 55	Leu	Ser	Cys	Leu	Thr 60	Val	Pro	Ala	Ser			
Ala 65	Val	Pro	Tyr	Arg	Asn 70	Ala	Ser	Gly	Ile	Tyr 75	His	Val	Thr	Asn	Asp 80			
Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Tyr	Glu	Ala 90	Asp	Asn	Leu	Ile	Leu 95	His			
Ala	Pro	Gly	Cys 100	Val	Pro	Cys	Val	Met 105	Thr	Gly	Asn	Val	Ser 110	Arg	Cys			
Trp	Val	Gln 115	Ile	Thr	Pro	Thr	Leu 120	Ser	Ala	Pro	Ser	Leu 125	Gly	Ala	Val			
Thr	Ala 130	Pro	Leu	Arg	Arg	Ala 135	Val	Asp	Tyr	Leu	Ala 140	Gly	Gly	Ala	Ala			
Leu 145	Cys	Ser	Ala	Leu	Tyr 150	Val	Gly	Asp	Ala	Cys 155	Gly	Ala	Leu	Phe	Leu 160			
Val	Gly	Gln	Met	Phe 165	Thr	Tyr	Arg	Pro	Arg 170	Gln	His	Ala	Thr	Val 175	Gln			
Asn	Cys	Asn	Cys 180	Ser	Ile	Tyr	Ser	Gly 185	His	Val	Thr	Gly	His 190	Arg	Met			
Ala																		
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cgt	cgcaa	agg g	getei	cgca	ac a	eggte	gtgag	g ggt	cctt	gag	gac	gggt	caa a	actat	gcaac	12	20	
agg	gaati	tta d	ccq	gttg	ct c	tttct	ctat	ctt	tatt	ctt	gcto	ettet	ct o	gtgt	ctgac	18	30	

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<210> 48

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 48

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Ile Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His 85 90 95

Ala Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys 100 105 110

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val 115 120 125

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala 130 135 140 Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 165 170 175

Asn Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met 180 185 190

Ala

<210> 49

<211> 959

<212> DNA

<213> hepatitis C virus

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<211> 319

<212> PRT

<213> hepatitis C virus

<400> 50

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro 100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 180 185 190

Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 195 200 205 Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 210 215 220

Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 225 230 235 240

Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 . 255

Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 265 270

Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 275 280 285

Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln Asn Cys 290 295 300

Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met Ala 305 310 315

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<211> 959

<212> DNA

<213> hepatitis C virus

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<210> 52

<211> 319

<212> PRT

<213> hepatitis C virus

<400> 52

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro 100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 170 165 Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 185 180 pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 205 200 195 Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 215 210 Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 235 230 225 Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 280 Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln Asn Cys 300 295 290 Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met Ala 310 305 <210> 53 <211> 959 <212> DNA <213> hepatitis C virus <400> 53 ccatgagcac gaatcctaaa cctcaaagaa aaaccaaaag aaacaccaac cgtcgcccac 60 aggacgtcaa gttcccgggc ggtggtcaga tcgttggcgg agtttacttg ttgccgcgca 120 ggggccctag gatgggtgtg cgcgcgactc ggaagacttc ggaacggtcg caaccccgtg 180 240 gacggcgtca gcctattccc aaggcgcgcc agcccacggg ccggtcctgg ggtcaacccg 300 ggtaccettg gcccetttac gccaatgagg gcctcgggtg ggcagggtgg ctgctctccc

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Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 180 · 185 190

Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 195 200 205

Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 210 215 220

Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 225 230 235 240

Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 255

Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 265 270

Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 275 280 285

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60

120

180

240

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Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly Ile Gly Thr Val Leu
                            40
Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Xaa Thr Xaa
                        55
Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu Glu Val
                    70
                                        75
65
Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr Xaa Arg Ala Ile Pro
                85
Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile Phe Cys His Ser Lys
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105

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gatgagatgg aggagtgctc ggcctcgttg ccctatatgg acgaaacacg tnccattgcc
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120 125

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Ile Ile Pro Asp Arg Glu Ala Leu Tyr Xaa Gln Phe Asp Glu Met Glu 50 55 60

Glu Cys Ser Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala 65 70 75 80

Gly Gln Phe Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln 85 90 95

Lys Ala Glu Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala 100 105 110

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Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Lys Ala 65 70 75 80

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Ala

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35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

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Ala

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<210> 111

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 111

Ser Thr Val Thr Glu Lys Asp Ile Arg Val Glu Glu Glu Val Tyr Gln

1 10 15

Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Thr Ala Leu Thr 20 25 30

Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

Ser Gly Leu Arg Asp Cys Thr Met Leu Val Tyr Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Ala Leu Gly
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Val

<210> 112

<211> 340

<212> DNA

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catgcataac agcaagggag acctgtgtgg gtatcgcaga tgccgcgcaa gcggcgtcta 180
caccaccagc ttcgggaaca cactgacgtg ctacctcaaa gcctcagccg ctatcagagc 240
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<210> 113

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 113

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Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Ala Ile Thr Ala Leu Thr 20 25 30

Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

Ala Gly Leu Arg Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Ala Leu Gly
100 105 110

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catgtacaac tccaaggggg acctatgcgg gcaacggagg tgccgcgcaa gcggggtcta	a
caccaccage ttegggaaca etgtaacgtg ttateteaag geegttgegg etaetaggge	2
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gageggegge gtagaggagg atgeaagage eeteegagee	
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Ser Thr Val Thr Glu Arg Asp Ile Arg Thr Glu Glu Glu Ile Tyr Gln 1 10 15	
Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Ser Ala Leu Thr 20 25 30	
Glu Arg Leu Tyr Val Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu 35 40 45	
Cys Gly Gln Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60	
Gly Asn Thr Val Thr Cys Tyr Leu Lys Ala Val Ala Ala Thr Arg Ala 65 70 75 80	
Ala Gly Leu Lys Gly Cys Ser Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95	

Val Ile Cys Glu Ser Gly Gly Val Glu Glu Asp Ala Arg Ala Leu Arg 

Ala

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gga	gccc	gag	gccc	gcaa	gg t	aata	gccg	c cc	tcac	ggag	aga	ctct	acg	tggg	cggccc
cat	gcat	aac	agca	aggg	ag a	cctt	tgcg	g gt	atcg	taga	tgc	cgcg	ıcga	gcgg	cgtata
cac	cacc	agc	ttcg	ggaa	ca c	aatg	acgt	g ct	acct	taag	gcc	tcag	cag	ccat	cagggc
tgc	<b>3</b> 999	cta	aagg	attg	ca c	catg	ctgg	t tt	gcgg	tgac	gac	ctag	tcg	tgat	cgccga
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Cys	Cys	Asp	Leu 20	Glu	Pro	Glu	Ala	Arg 25	Lys	Val	Ile	Ala	Ala 30	Leu	Thr
Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	His	Asn	Ser	Lys 45	Gly	Asp	.Leu
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Tyr 60	Thr	Thr	Ser	Phe
Gly 65	Asn	Thr	Met	Thr	Cys 70	Tyr	Leu	Lys	Ala	Ser 75	Ala	Ala	Ile	Arg	Ala 80
Ala	Gly	Leu	Lys	Asp 85	Cys	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val
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gggaatcttc ccggttgctc tttctctatc ttcctcttgg cacttctt	tc gtgcctgact										
gttcccacct cggccgtcaa ctatcgcaat gcctcgggca tctatcaca	at caccaatgac										
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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val A 20 25 3	Arg Ala Val 30										
Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly C 35 40 45	Cys Ser Phe										
Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val P 50 55 60	Pro Thr Ser										
Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile T 65 70 75	Thr Asn Asp 80										

Cys Pro Asn Ser Ser Ile Val Tyr Glu Thr Glu His His Ile Leu His

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Trp Val Ala Leu Thr Pro Thr Val Ala Ala Pro Tyr Ile Gly Ala Pro 115 120 125

Leu Glu Ser Leu Arg Ser His Val Asp Leu Met Val Gly Ala Ala Thr
130 135 140

Ala Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Val Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Val Thr Gly His Arg 180 185 190

<210> 120

<211> 574

<212> DNA

<213> hepatitis C virus

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<210> 121

<211> 191

<212> PRT

<213> hepatitis C virus

<400> 121

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Gln His Tyr Arg Asn Ile Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Thr Ser Arg Cys
100 105 110

Trp Val Pro Leu Thr Pro Thr Val Ala Ala Pro Tyr Val Gly Ala Pro
115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr
130 135 140

Val Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Arg Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Asp Gly His Ile Thr Gly His Arg 180 185 190

<210> 122

<211> 574

<212> DNA

<213> hepatitis C virus

<400> 122

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gggaatcttc	ccggttgctc	tttctctatc	ttcctcctgg	cacttctttc	gtgcctcact	180
gtcccagcgt	cagctgagca	ctaccggaat	gcttcgggca	tctatcacat	caccaatgac	240
tgtccgaatt	ccagcgtagt	ctatgaaact	gaccaccata	tattgcactt	gccggggtgc	300
gtaccctgcg	tgagggccgg	gaacgtgtct	cgttgctgga	cgccggtaac	acctacggtg	360
gctgccgtat	ccatggacgc	tccgctcgag	tccttccggc	ggcatgtgga	cctaatggta	420
ggtgcggcca	ccgtgtgttc	tgtcctctat	gttggagacc	tctgtggagg	tgctttccta	480
gtggggcaga	tgttcacctt	ccagccgcgt	cgccactgga	ccacgcagga	ttgtaattgc	540
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<210> 123

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<213> hepatitis C virus

<400> 123

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Glu His Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Val Val Tyr Glu Thr Asp His His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Ala Gly Asn Val Ser Arg Cys
100 105 110

Trp Thr Pro Val Thr Pro Thr Val Ala Ala Val Ser Met Asp Ala Pro 115 120 125

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Leu Glu Ser Phe Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr
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  145
                      150
  Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln
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                  165
  Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg
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Glu His Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile
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Val Tyr Glu Thr Glu His His Ile Leu His Leu
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Val Tyr Glu Ala Asp His His Ile Met His Leu
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Val Tyr Glu Thr Asp His His Ile Leu His Leu
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Val Arg Thr Gly Asn Thr Ser Arg Cys Trp Val Pro Leu
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<212> PRT
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Val Arg Ala Gly Asn Val Ser Arg Cys Trp Thr Pro Val
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Ala Pro Tyr Ile Gly Ala Pro Leu Glu Ser
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Ala Pro Tyr Val Gly Ala Pro Leu Glu Ser
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<212> PRT

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Ala Val Ser Met Asp Ala Pro Leu Glu Ser
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Gln Pro Arg Arg His Trp Thr Thr Gln Asp
               5
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Arg Pro Arg Arg His Trp Thr Thr Gln Asp
             5
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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Asp Arg Pro Thr Gly Lys Ser Trp Gly Lys Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Val Pro Val Ser Gly Leu 180 185 190

Gln Val Lys Asn Thr Ser Ser Ser Tyr Met Val Thr Asn Asp Cys Gln
195 200 205

Asn Ser Ser Ile Val Trp Gln Leu Arg Asp Ala Val Leu His Val Pro 210 215 220

Gly Cys Val Pro Cys Glu Glu Lys Gly Asn Ile Ser Arg Cys Trp Ile

Pro Val Ser Pro Asn Ile Ala Val Ser Gln Pro Gly Ala Leu Thr Lys 255 245 Gly Leu Arg Thr His Ile Asp Thr Ile Ile Ala Ser Ala Thr Phe Cys 265 Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Ala Val Met Leu Ala Ser 280 Gln Val Phe Ile Ile Ser Pro Gln His His Lys Phe Val Gln Asp Cys 295 290 Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala 315 305 310 <210> 145 340 <211> <212> DNA <213> hepatitis C virus <400> 145 ctcaacggtc acggagaggg acatcagaac tgaggagtcc atataccttg cttgctcttt 60 accogagoag goacggactg coatacacto actgactgag aggotttacg tgggagggco 120 catgctaaac agcaaagggc aaacctgcgg atacagacgc tgccgcgcca gcggagtgtt 180 caccactage atgggaaata ccatcacgtg ctacgtgaag gcacaagcag cctgtaagge tgcgggcata attgccccca cgatgctggt gtgcggcgac gatctagttg tcatctcaga 300 340 gagtcagggg accgaggagg acgagcggaa cctacgagcc <210> 146 <211> 113 <212> PRT

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<400> 146

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Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	Leu	Asn	Ser	Lys 45	Gly	Gln	Thr	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Phe 60	Thr	Thr	Ser	Met	
Gly 65	Asn	Thr	Ile	Thr	Cys 70	Tyr	Val	Lys	Ala	Gln 75	Ala	Ala	Суз	Lys	Ala 80	
Ala	Gly	Ile	Ile	Ala 85	Pro	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
Val	Ile	Ser	Glu 100	Ser	Gln	Gly	Thr	Glu 105	Glu	Asp	Glu	Arg	Asn 110	Leu	Arg	
Ala																
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acco	ttgg	jee e	ctat	atgg	g aa	tgag	ggct	gcg	ggtg	ggc	aggg	tggc	tc c	tgtc	cccgc	300
gcgg	gctct	cg c	ccgt	cgtg	ia aa	ccca	aatg	acc	cccg	gcg	cagg	а				345
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Pro	Gly	His	Arg 20	Thr	Leu	Ser		Gln 25	Ala	Ala	Val		Ser 30	Leu	Val	

Glu Phe Thr Cys Tyr His Ala Gly Ala Pro Ser Trp Val Cys Val Gln 40 Cys Ala Arg Leu Pro Ser Gly Arg Asn Leu Ala Val Gly Ala Asn Pro 55 Ser Pro Gly Arg Ala Glu Pro Arg Ala Gly Pro Gly Leu Ser Pro Gly 75 70 Thr Leu Gly Pro Tyr Met Gly Met Arg Ala Ala Gly Gly Gln Gly Gly 90 Ser Cys Pro Arg Ala Ala Leu Ala Arg Arg Gly Ala Gln Met Thr Pro 100 Gly Ala Gly 115 <210> 149 280 <211> <212> DNA <213> hepatitis C virus <400> 149 ggcctgtgac ctcaaggacg aggctaggag ggtgataact tcactcacgg agcggcttta 60 120 ctgtggtggt cctatgttca acagcaaggg acaacactgc ggttaccgcc gctgccgtgc tagtggggtg ctacccacca gcttcgggaa cacaatcacc tgttacatca aagcaaaggc 180 agctaccaaa gctgccggaa ttaaaaatcc atcattcctt gtctgcggag atgacttggt 240 280 cgtgattgct gagagtgcag ggatcgatga ggacagagcg <210> 150 <211> 93 <212> PRT <213> hepatitis C virus <400> 150 Ala Cys Asp Leu Lys Asp Glu Ala Arg Arg Val Ile Thr Ser Leu Thr

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20

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Pro Thr Ser Phe 40 35 Gly Asn Thr Ile Thr Cys Tyr Ile Lys Ala Lys Ala Ala Thr Lys Ala 50 Ala Gly Ile Lys Asn Pro Ser Phe Leu Val Cys Gly Asp Asp Leu Val Val Ile Ala Glu Ser Ala Gly Ile Asp Glu Asp Arg Ala <210> 151 <211> 499 <212> DNA <213> hepatitis C virus <400> 151 atgagcacga atcctaaacc tcaaagaaaa accaaaagaa acaccaaccg tcgcccacag 60 gacgtcaagt tcccgggcgg tggtcagatc gttggcggag tttacttgtt gccgcgcagg 120 180 qqccctaqqa tgggtgtgcg cgcgactcgg aagacttcgg aacggtcgca accccgtgga 240 cqqcqtcaqc ctattcccaa ggcgcgccag cccacgggcc ggtcctgggg tcaacccggg 300 taccettqqc ccctttacgc caatgagggc ctcgggtggg cagggtggct gctctcccct cqaqqctctc ggcctaattg gggccccaat gacccccggc gaaaatcgcg taatttgggt 360 420 aaggtcatcg ataccctaac gtgcggattc gccgatctca tggggtatat cccgctcgta qqcqqcccca ttgggggcgt cgcaagggct ctcgcacacg gtgtgagggt ccttgaggac 480 499 ggggtaaact atgcaacag <210> 152 <211> 166 <212> PRT <213> hepatitis C virus

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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala

35

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<400> 154

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Glu Asp Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 55 50

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 75 70 65

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His 95 90 85

Ala Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val 115 120

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala 130 135

Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 155 160 145 150

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Asn Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met

180 185 190

Ala

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<213> hepatitis C virus

<400> 156

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Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

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Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 75	
/5 ASP	
Cys Pro Asn Ser Ser 73	
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Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Leu Ile Leu His	
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Ala Pro Gly Cys Val Pro Cys Val Arg Lys Asp Asn Val Ser Arg Cys  100 105	
100 Cys Val Arg Lys Asp Asp	
105 ASN Val Ser Arg Cvs	
Tro. 11.0	
of Val Gln Ile Thr Pro The .	
Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Phe Gly Ala Val	
120 Phe Gly Ala Val	
Thr Ala P	
130 Leu Arg Arg Ala Val	
135 Asp Tyr Leu Val Gly Gl	
Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Val Gly Gly Ala Ala 130 135 140	
Leu Cvs Ser al	
Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 150 155	
150 Asp Ala Cys Gly Ala Leu Phase	
155 Leu	
Val Gly Gln Met Di	
The Thr Tyr Arg Pro Are G	
Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 170	
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Ala Tyr Tyr Ala Ser Val Ala Asn Trp Thr Lys Val Val Leu Val Leu

Phe Leu Phe Ala Gly Val Asp Ala Thr Thr Gln Ile Ser Gly Gly Ser

Ser Ala Gln Thr Thr Tyr Gly Ile Ala Ser Phe Ile Thr Arg Gly Ala

Gln Gln Lys Leu Gln Leu Ile Asn Thr Asn Gly Ser Trp His Ile Asn

Arg Thr Ala Leu Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Ile Ala

Gly Leu Phe Tyr Tyr His Lys Phe Asn Ser Ser Gly Cys Pro Asp Arg

Met Ala Ser Cys Arg Ala Leu Ala Thr Phe Asp Gln Gly Trp Gly Thr

Ile Ser Tyr Ala Asn Ile Ser Gly Pro Ser Asp Asp Lys Pro Tyr Cys

Trp His Tyr Pro Pro Arg Pro Cys Gly Val Val Pro Ala Gln Glu Val <210> 159

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Gln Arg Leu Tyr Cys Gly Gly Pro Met Tyr Asn Ser Lys Gly Gln Gln
35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Met 50 55 60

Gly Asn Thr Met Thr Cys Tyr Ile Lys Ala Leu Ala Ser Cys Arg Ala 65 70 75 80

Ala Arg Leu Arg Asp Cys Thr Leu Leu Val Cys Gly Asp Asp Leu Val 85 90 95

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Ala .

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<210> 162 <211> 113 <212> PRT <213> hepatitis C virus

<400> 162

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Ser Cys Asp Ser Gln Pro Asp Ala Arg Ala Ala Ile Arg Ser Leu Thr 20 25 30

Gln Arg Leu Phe Cys Gly Gly Pro Met Tyr Asn Ser Lys Gly Gln Gln 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Met 50 55 60

Gly Asn Thr Met Thr Cys Tyr Ile Lys Ala Leu Ala Ser Cys Arg Thr 65 70 75 80

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Ala Ile Cys Glu Ser Gln Gly Thr His Glu Asp Glu Ala Asn Leu Arg 100 105 110

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120

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240

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480

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90

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Val Gly Gly Ala Ala 130 Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 145 Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 165 Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Gln Met 185 180 Ala <210> 197 <211> 1485 <212> DNA <213> hepatitis C virus <220> <221> misc\_feature <222> (787)..(787) <223> "n" is any nucleotide <220> <221> misc\_feature <222> (1011)..(1011) <223> "n" is any nucleotide <220> <221> misc\_feature <222> (1444)..(1444) <223> "n" is any nucleotide <400> 197 tgtgccagga ccatcaccac cggagcttct atcacatact ccacttacgg caagttcctt 60 gctgatggag ggtgttcagg cggcgcgcat gacgtgatca tatgcgacga gtgccattcc 120 caggacgcca ccaccattct tgggataggc actgtccttg accaggcaga gacggctgga 180 getaggeteg tegtettgge caeggecaec ceteceggea gtgtgacaac geeceaecee 240 aacatcgagg aagtggccct gcctcaggag ggggaggttc ccttctacgg cagagccatt 300

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Ile Ile Cys Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly 35 40 45

Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val 50 . 55 60

Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro 65 70 75 80

Asn Ile Glu Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr 85 90 95

Gly Arg Ala Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile 100 105 110

Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr 115 120 125

Ser Leu Gly Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala 130 135 140

Val Ile Pro Thr Thr Gly Asp Val Val Val Cys Ser Thr Asp Ala Leu 145 150 155 160

Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser 165 170 175

Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile 180 185 190 Glu Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Trp
195 200 205

Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala 210 215 220

Gly Glu Arg Pro Ser Gly Met Phe Asp Ser Val Val Leu Cys Glu Cys 225 230 235 240

Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr 245 250 255

Val Arg Leu Arg Ala Tyr Xaa Asn Thr Pro Gly Leu Pro Val Cys Gln 260 265 270

Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile 275 280 285

Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Glu Asn Phe Pro 290 295 300

Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro 305 310 315

Pro Pro Ser Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro 325 330 335

Xaa Leu Thr Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln 340 345 350

Asn Glu Ile Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys 355 360 365

Met Ser Ala Asp Leu Glu Val Ile Thr Ser Thr Trp Val Leu Val Gly 370 375 380

Gly Val Val Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val 385 390 395 400

Ala Ile Val Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro 405 410 415

Asp Arg Glu Val Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser

420 425 430

Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe 440 435 Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu 455 Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe 475 465 470 Trp Xaa Thr Tyr <210> 199 <211> 1485 <212> DNA <213> hepatitis C virus <220> <221> misc\_feature <222> (684)..(684) <223> "n" is any nucleotide <220> <221> misc feature <222> (909)..(909) <223> "n" is any nucleotide <220> <221> misc\_feature <222> (1058)..(1058) <223> "n" is any nucleotide <220> <221> misc\_feature <222> (1133)..(1133) "n" is any nucleotide <223> <400> 199 tgtgccagga ccatcaccac cggagcttct atcacatact ccacttacgg caagttcctt 60 gctgatggag ggtgttcagg cggcgcgtat gacgtgatca tatgcgacga gtgccattcc 120 caggacgcca ccaccattct tgggataggc actgtccttg accaggcaga gacggctgga 180

gctaggctcg tcgtcttggc cacggccacc cctcccggca gtgtgacaac gccccacccc

300 aacatcgagg aagtggccct gcctcaggag ggggaggttc ccttctacgg cagagccatt ccccttgctt ttataaaggg tggtaggcat ctcatcttct gccattccaa gaaaaaatgt 360 gatgaactcg ccaagcaact gaccagcctg ggcgtgaacg ccgtggcata ttatagaggt 420 480 ctagacgtcg ccgtcatccc cacagcagga gacgtggtcg tgtgcagcac cgacgcgctc atgacgggat tcaccggcga ctttgattct gtcatagact gcaactccgc cgtcactcag 540 acggtggact tcagtctgga tcccactttt accattgaga ctaccacagt gccccaggac 600 gcagtgtcca gaagccagcg taggggccgc acggggagag gtaggcacgg catataccgg 660 tatgtctcgg ctggagagag accntctgac atgttcgact ccgtggtgct ctgtgagtgc 720 tacgatgccg gatgtgcgtg gtatgatctg actectgccg agactaccgt gaggttgcgc 780 840 gettacataa acaccccgg getecetgte tgtcaggace atttggaatt etgggagggg gtgttcacgg ggctcactaa catcgacgct cacatgctgt cacagaccaa acagggtggg 900 gagaatttnc cataccttgt agcgtaccaa gcaacagtct gtgttcgcgc gaaagcgccc 960 cccccagct gggacacaat gtggaaatgc atgctccgtc tcaaaccgac tttaactggc 1020 cctactcccc tcttgtacag gctggggccc gtccagantg agatcacact gacgcacccc 1080 1140 atcaccaagt acattatggc ttgcatgtct gcggacttgg aggtcattac cancacttgg 1200 gttctggtgg ggggcgttgt ggcggccctg gcggcctact gcttgacggt gggttcggta gccatagtcg gtaggatcat cctctctggg aaacctgcca tcattcccga tagggaggca 1260 1320 ttataccage aatttgatga gatggaggag tgctcggcct cgttgcccta tatggacgag 1380 acacgtgcca ttgccggaca attcaaagag aaagtgctcg gcttcatcag cacgaccggc cagaaggctg aaactctgaa gccggcagcc acgtctgtgt ggaacaaggc tgagcagttc 1440 1485 tgggccacat acatgtggaa cttcatcagc gggatacaat aatag

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<210> 200
<211> 484
<212> PRT
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<sup>&</sup>lt;213> hepatitis C virus

<sup>&</sup>lt;220>

<sup>&</sup>lt;221> MISC\_FEATURE

<sup>&</sup>lt;222> (228)..(228)

<sup>&</sup>lt;223> "Xaa" is any amino acid

<sup>&</sup>lt;220>

<sup>&</sup>lt;221> MISC\_FEATURE

<sup>&</sup>lt;222> (303)..(303)

<223> "Xaa" is any amino acid

<220>

<221> MISC\_FEATURE

<222> (353)..(353)

<223> "Xaa" is any amino acid

<220>

<221> MISC\_FEATURE

<222> (378)..(378)

<223> "Xaa" is any amino acid

<400> 200

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Gly Lys Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Val 20 25 30

Ile Ile Cys Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly 35 40 45

Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val 50 55 60

Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro 65 70 75 80

Asn Ile Glu Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr 85 90 95

Gly Arg Ala Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile 100 105 110

Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr 115 120 125

Ser Leu Gly Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala 130 135 140

Val Ile Pro Thr Ala Gly Asp Val Val Cys Ser Thr Asp Ala Leu 145 150 155 160 Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser 165 170 175

Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile 180 185 190

Glu Thr Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Arg 195 200 205

Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala 210 215 220

Gly Glu Arg Xaa Ser Asp Met Phe Asp Ser Val Val Leu Cys Glu Cys 225 230 235 240

Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr 245 250 255

Val Arg Leu Arg Ala Tyr Ile Asn Thr Pro Gly Leu Pro Val Cys Gln
260 265 270

Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile 275 280 285

Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Glu Asn Xaa Pro 290 295 300

Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro 305 310 315 320

Pro Pro Ser Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro 325 330 335

Thr Leu Thr Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln 340 345 350

Xaa Glu Ile Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys 355 360 365

Met Ser Ala Asp Leu Glu Val Ile Thr Xaa Thr Trp Val Leu Val Gly 370 380

Gly Val Val Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val

385	390	395	400

Ala Ile Val Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro 405 410 415

Asp Arg Glu Ala Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser 420 425 430

Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe 435 440 445

Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu 450 455 460

Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe 465 470 475 480

Trp Ala Thr Tyr

<210> 201

<211> 340

<212> DNA

<213> hepatitis C virus

<400> 201

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<210> 202

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 202

Ser Thr Val Thr Glu Arg Asp Ile Arg Val Glu Glu Glu Val Tyr Gln 1 5 10 15

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 120 115. Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val 130 Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp Gly Ile Asn Tyr Ala Thr <210> 165 499 <211> <212> DNA hepatitis C virus <213> <220> <221> misc\_feature (306) . . (306) <222> <223> "n" is any nucleotide <220> misc feature <221> (313)..(313) <222> "n" is any nucleotide <223> <220> <221> misc\_feature (340)..(340) <222> "n" is any nucleotide <223> atgagcacga atcctaaacc tcaaagaaaa accaaacgta acaccaaccg ccgccctatg 60 120 gacgttaagt tcccaggcgg tggtcagatc gttggcggag tttacttgtt gccgcgcagg ggccccaggt tgggtgtgcg cgcgactcgg aagacttcgg agcggtcgca acctcgtggg

aggcgccaac ctatcccaa ggcgcgccga accgagggca gatcctggg gcagcccggg 240
tatccttggc ccctttacgg caatgagggc tgtgggtggg cagggtggct cctgtccct 300
cgcggntctc ggncgtcttg gggccccaat gatcccggn ggagatcccg caacttgggt 360
aaggtcatcg ataccctaac atgcggcttc gccgacctca tgggatacat cccgcttgta 420

Α.

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gggatcaatt atgcaacag . 499

<210> 166

<211> 126

<212> PRT

<213> hepatitis C virus

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<223> "Xaa" is any amino acid

<220>

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<222> (105)..(105)

<223> "Xaa" is any amino acid

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<221> MISC\_FEATURE

<222> (114)..(114)

<223> "Xaa" is any amino acid

<400> 166

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Arg Thr Glu Gly Arg Ser Trp Ala Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Xaa Ser Arg Xaa Ser Trp Gly Pro Asn Asp Pro 100 105 110

Arg Xaa Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu 115 120 125

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<210> 168 <211> 193 <212> PRT

<213> hepatitis C virus

<400> 168

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser 50 55 60

Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Glu Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Ala Gly Ala Pro 115 120 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr
130 135 140

Met Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg Met 180 185 190

Ala

<210> 169

<211> 579

<212> DNA

<213> hepatitis C virus

<400> 169

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<211> 193

<212> PRT

<213> hepatitis C virus

<400> 170

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Gly Val Asn Tyr Arg Asn Ala Ser Gly Val Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Asp Asn His Ile Leu His
85 90 95

Leu Pro Gly Cys Val Pro Cys Val Lys Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Val Gly Ala Pro 115 120 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Val Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Ile Thr Gly His Arg Met 180 185 190 <210>

171 579

<211> <212> DNA hepatitis C virus <400> 171 acatgcggct tcgccgacct catgggatac atcccgcttg tgggcgcccc tgttggtggc gtcgccagag cccttgcgca cggcgtcagg gctgtggaag acgggattaa ctatgcaaca qqqaaccttc ctggttgctc cttttctatc ttccttctgg cacttctctc gtgcctgact gtccccgcct cggctgtgca ttatcacaac acctcgggca tctaccacct caccaatgac tgccctaact ctagcatagt ctttgaggca gtccatcaca tcttgcacct tccaggatgc gtcccttgtg taagaactgg gaaccagtct cggtgctggg tagccttgac ccccacgctg qccqcqccat accttgqcgc tccactcgag tccatgcggc gtcacgtgga tttgatggtg ggcactgcta cattgtgctc agcactctac gttggggacc tgtgcggggg catattccta gegggccaga tgttcacctt ceggeceege etecattgga ecaeceagga gtgcaattgt tccacctatc cgggccacat cacgggtcat agaatggcg <210> 172 <211> 193 <212> PRT <213> hepatitis C virus <400> 172 Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 25 20 Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55

Ala Val His Tyr His Asn Thr Ser Gly Ile Tyr His Leu Thr Asn Asp

60

120

180

240

300

360

420

480

540

65 70 75 80

Cys Pro Asn Ser Ser Ile Val Phe Glu Ala Val His His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Leu Ala Ala Pro Tyr Leu Gly Ala Pro
115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Thr Ala Thr
130 135 140

Leu Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Ile Phe Leu 145 150 155 160

Ala Gly Gln Met Phe Thr Phe Arg Pro Arg Leu His Trp Thr Thr Gln 165 170 175

Glu Cys Asn Cys Ser Thr Tyr Pro Gly His Ile Thr Gly His Arg Met 180 185 190

Ala ·

<210> 173

<211> 579

<212> DNA

<213> hepatitis C virus

<400> 173

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<210> 174

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 174

Thr Cys Gly Ser Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val His Tyr His Asn Thr Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Phe Glu Ala Glu His His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Ile Ala Leu Thr Pro Thr Leu Ala Ala Pro His Ile Gly Ala Pro 115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Thr Ala Thr 130 135 140

Leu Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Ile Phe Leu 145 150 155 160

Val Gly Gln Met Phe Asn Phe Arg Pro Arg Leu His Trp Thr Thr Gln 165 170 175

Glu Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met

Ala

<210><211><211>	175 579 DNA			·			
		atitis C vi	rus				
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gggaato	ettc	ccggttgctc	cttttctatc	ttccttctag	cacttctctc	gtgcttgact	180
gtcccgg	gcct	cggcgcagca	ctaccggaac	atctcgggca	tttatcacgt	caccaatgac	240
tgcccga	act	ctagtatagt	gtatgaagct	gaccatcata	tcatgcatct	accagggtgt	300
gtgcctt	gcg	tgagaaccgg	gaacacctcg	cgctgctggg	ttcctttaac	acccactgtg	360
gctgcc	cct	atgttggcgc	gccgctcgaa	tccatgcggc	ggcacgtgga	cttaatggtg	420
ggtgccg	gcca	ccgtctgctc	ggccctgtac	atcggagacc	tttgcggagg	tgtcttcctg	480
gtcgggc	caga	tgttcacctt	ccggccgcgc	cgccattgga	ctacccagga	ctgcaactgc	540
tctatct	atg	atggccacat	caccggccat	agaatggct			579
<210>	176			•			

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 176

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 5 10

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 30 20

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 55

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Thr Ser Arg Cys 100 105 Trp Val Pro Leu Thr Pro Thr Val Ala Pro Tyr Val Gly Ala Pro Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr Val Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 150 155 145 Val Gly Gln Met Phe Thr Phe Arg Pro Arg Arg His Trp Thr Thr Gln 165 170 Asp Cys Asn Cys Ser Ile Tyr Asp Gly His Ile Thr Gly His Arg Met Ala <210> 177 <211> 579 <212> DNA <213> hepatitis C virus <400> 177 acgtgcgggt tcgccgacct catgggatac atcccgctcg tgggcgctcc agtaggaggc 60 gtcgccagag ccttggcgca tggcgtcagg gctgtggagg acgggatcaa ttacgcaaca gggaacette ceggetgete ettitetate tiectetigg tactietete gegeetaact 180 gtcccagcgt ctgctcagca ctaccggaat gcatcgggca tctaccatgt caccaacgac 240 tgcccgaact ccagtattgt gtatgaagcc gaccatcaca tcatgcacct acccgggtgt 300

gtgccctgtg taagaactgg gaatgtctcg cgttgctgga ttcctttaac acccactgta

geogtecect acetegggge tecaettacg tetgtacgge ageatgtgga cetgatggtg

ggggcggcca ccttatgctc tgccctctac atcggagacc attgcggagg tgtcttcttg

Ala Gln His Tyr Arg Asn Ile Ser Gly Ile Tyr His Val Thr Asn Asp

75

70 ·

360

420

<210> 178

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 178

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Val Leu Leu Ser Arg Leu Thr Val Pro Ala Ser 50 55 60

Ala Gln His Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Val Ser Arg Cys
100 105 110

Trp Ile Pro Leu Thr Pro Thr Val Ala Val Pro Tyr Leu Gly Ala Pro 115 120 125

Leu Thr Ser Val Arg Gln His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Leu Cys Ser Ala Leu Tyr Ile Gly Asp His Cys Gly Gly Val Phe Leu 145 150 155 160

Ala Gly Gln Met Val Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

## Asp Cys Asn Cys Ser Ile Tyr Val Gly His Ile Thr Gly His Arg Met 180 185 190

Ala

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<210> 179
<211> 579
<212> DNA
<213> hepatitis C virus
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<221> misc_feature
<222> (568)..(579)
<223> "n" is any nucleotide
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gtcgccagar ctctggcgca tggcgtcagg gctctggaag acgggatcaa ttatgcaaca
                                                                     120
                                                                     180
qqqaatcttc ctggttgctc tttctctatc tcccttcttg aacttctctc gtgcctgact
gttcccgcct cagccatcca ctatcgcaat gcttcggacg gttattatat caccaatgat
                                                                     240
tgcccqaact ctaqcataqt gtatgaagcc gagaaccaca tcttgcacct tccggggtgt
                                                                     300
ataccetgtg tgaagaccgg gaatcagteg cggtgetggg tggeteteac ecceaegetg
                                                                     360
geggeeecae acetaegtge teegettteg teettaeggg egeatgtgga eetaatggtg
                                                                     420
ggggccgcca cggcatgctc cgctttttac attggagatc tgtgcggggg tgtgtttttg
                                                                     480
gegggecaac tgttcactat eeggecaege atteatgaaa eeacteagga etgeaattge
                                                                     540
                                                                     579
tccatctact cagggcacat cacgggtnnn nnnnnnnn
<210> 180
<211> 193
<212> PRT
<213> hepatitis C virus
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<221> MISC_FEATURE
<222> (24)..(24)
<223> "Xaa" is any amino acid
<220>
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<222> (190)..(193)
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<223> "Xaa" is any amino acid

<400> 180

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Xaa Leu Ala His Gly Val Arg Ala Leu 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Ser Leu Leu Glu Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Ile His Tyr Arg Asn Ala Ser Asp Gly Tyr Tyr Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Glu Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Ile Pro Cys Val Lys Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Leu Ala Ala Pro His Leu Arg Ala Pro 115 120 125

Leu Ser Ser Leu Arg Ala His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Ala Cys Ser Ala Phe Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Ala Gly Gln Leu Phe Thr Ile Arg Pro Arg Ile His Glu Thr Thr Gln 165 170 175

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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Xaa Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser

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Trp	Val	Ser 115	Leu	Thr	Pro	Thr	Val 120	Ala	Ala	Pro	Tyr	Leu 125	Gly	Ala	Pro	
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Val	Gly	Gln	Met	Phe 165	Thr	Phe	Gln	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln	
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Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln

Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg Met 180 185

Ala

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<210> 186

<211> 193 <212> PRT

<213> hepatitis C virus

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Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 40 45 35

Ser Ile Tyr Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser • 55 Ala Ile His Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp 70 Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Leu His 90 85 Leu Pro Gly Cys Leu Pro Cys Val Arg Val Gly Asn Gln Ser Arg Cys Trp Val Ala Leu Ser Pro Thr Val Ala Ala Pro Tyr Ile Gly Ala Pro 115 120 Val Glu Ser Phe Arg Arg His Val Asp Met Met Val Gly Ala Ala Thr 135 Val Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 150 145 Val Gly Gln Met Phe Ser Phe Arg Pro Arg Arg His Trp Thr Thr Gln 165 Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Ile Thr Gly His Gly Met 180 Ala <210> 187 <211> 579 <212> DNA <213> hepatitis C virus <400> 187 acttgcggct ttgccgacct catgggatac atcccgctcg taggcgcccc tgtgggtggc 60 qtcgccaggg ccctggcaca cggtgttagg gctgtggagg acgggatcaa ttatgcgaca 120 gggaatette eeggttgete titetetate tieetetigg eactietite gigeetgaet 180 gttcccacct cggccgtcaa ctatcgcaat gcctcgggca tctatcacat caccaatgac 240 tgcccgaact cgagcatagt gtacgagacc gagcaccaca tcctacacct cccagggtgt 300 ttaccctgcg tgagggttgg gaatcagtca cgctgctggg tggccctcac tcccaccgtg 360

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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30	
Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45	
Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser 50 55 60	
Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80	
Cys Pro Asn Ser Ser Ile Val Tyr Glu Thr Glu His His Ile Leu His 85 90 95	
Leu Pro Gly Cys Leu Pro Cys Val Arg Val Gly Asn Gln Ser Arg Cys 100 105 110	
Trp Val Ala Leu Thr Pro Thr Val Ala Ala Pro Tyr Ile Gly Ala Pro 115 120 125	
Leu Glu Ser Leu Arg Ser His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140	
Ala Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160	

Val Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Val Thr Gly His Arg Met 180 185 190

Ala

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3343433		J	3 23 2		_	
gtggggcaga	tgttcacctt	ccagccgcgt	cgccactgga	ccacgcagga	ttgtaattgc	540
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<210> 190

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 190

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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe

Ser	Ile 50	Phe	Leu	Leu	Ala	Leu 55	Leu	Ser	Cys	Leu	Thr 60	Val	Pro	Ala	Ser		
Ala 65	Glu	His	Tyr	Arg	Asn 70	Ala	Ser	Gly	Ile	Tyr 75	His	Ile	Thr	Asn	Asp 80		
Cys	Pro	Asn	Ser	Ser 85	Val	Val	Tyr	Glu	Thr 90	Asp	His	His	Ile	Leu 95	His		
Leu	Pro	Gly	Cys 100	Val	Pro	Cys	Val	Arg 105	Ala	Gly	Asn	Val	Ser 110	Arg	Cys		
Trp	Thr	Pro 115	Val	Thr	Pro	Thr	Val 120	Ala	Ala	Val	Ser	Met 125	Asp	Ala	Pro		
Leu	Glu 130	Ser	Phe	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr		
Val 145	Cys	Ser	Val	Leu	Tyr 150	Val	Gly	Asp	Leu	Cys 155	Gly	Gly	Ala	Phe	Leu 160		
Val	Gly	Gln	Met	Phe 165		Phe	Gl'n	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln		
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Ala																	
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<211> 166

<212> PRT

<213> hepatitis C virus

<400> 194

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Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Ala Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp 145 150 155 160

Gly Ile Asn Tyr Arg Gln 165

<210> 195

<211> 579

<212> DNA

<213> hepatitis C virus

<400> 195

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<211> 193

<212> PRT

<213> hepatitis C virus

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Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Leu Ile Leu His 85 90 95

Ala Pro Gly Cys Val Pro Cys Val Arg Lys Asp Asn Val Ser Arg Cys
100 105 110

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Phe Gly Ala Val

Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Thr Ala Leu Thr 20 25 30	
Glu Arg Leu Tyr Val Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu 35 40 45	
Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60	
Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80	
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caccacaagc ttcgggaaca ctctgacgtg cttcctcaag gccacagcgg ccaccaaagc	240 300
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<210> 206

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 206

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1 5 10 15

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<210> 208

<211> 113

<212> PRT

<213> hepatitis C virus

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<400> 208

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Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Ser Ala Leu Thr

Glu Arg Leu Tyr Lys Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu

Cys Gly Leu Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe

Gly Asn Thr Val Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val

Val Ile Ala Glu Ser Gly Gly Val Glu Glu Asp Ala Arg Ala Leu Arg Ala

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Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 40

Cys Gly Ile Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 55

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala 70

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val

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Thr

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<213> hepatitis C virus

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<222> (327)..(327)

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<222> (340)..(340)

<223> "n" is any nucleotide

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catqttcaac agcaagggag acctgtgcgg acaacgccgg tgccgcgcaa gcggcgtgtt
caccaccage ttegggaaca caetgaegtg ctacettaaa gecacagetg ctaetagage
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gagegeeggt gtggaggagg atcccanaac cennegacen
<210> 212
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<211>
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<213> hepatitis C virus
<220>
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<222> (4)..(4)
<223> "Xaa" is any amino acid
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<222> (109)..(109)
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<222> (111) ...(111)
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Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Thr Ala Leu Thr
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            20
                                                    30
Glu Arq Leu Tyr Val Gly Gly Pro Met Phe Asn Ser Lys Gly Asp Leu
        35
                                                45
Cys Gly Gln Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Phe
    50
                        55
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Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala

60

120

180

240

300

70

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Ser Glu Ser Ala Gly Val Glu Glu Asp Pro Xaa Thr Xaa Arg 100 105 110

Pro

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cctgactaat tcaaaggggc agaactgtgg ctatcgccga tgccgcgcaa gcggcgtgct 180

gacgaccagc tgcggtaata cccttacatg ttacctaaag gcctctgcag cctgtcgagc 240

tgcgaagctc caggactgca cgatgctcgt gtgcggggac gaccttgtcg ttatctgtga 300

aagcgcggga acccaagagg acgcggcgag cctacgagtc 340

<210> 214 <211> 113 <212> PRT <213> hepatitis C virus

<400> 214

Ser Thr Val Thr Glu Asn Asp Ile Arg Val Glu Glu Ser Ile Tyr Gln
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Cys Cys Asp Leu Ala Pro Glu Ala Arg Gln Ala Ile Lys Ser Leu Thr 20 25 30

Glu Arg Leu Tyr Ile Gly Gly Pro Leu Thr Asn Ser Lys Gly Gln Asn 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Thr Thr Ser Cys 50 60

Gly 65	Asn	Thr	Leu	Thr	Cys 70	Tyr	Leu	Lys	Ala	Ser 75	Ala	Ala	Cys	Arg	Ala 80	
Ala	Lys	Leu	Gln	Asp 85	Cys	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
Val	Ile	Cys	Glu 100	Ser	Ala	Gly	Thr	Gln 105	Glu	Asp	Ala	Ala	Ser 110	Leu	Arg	
Val																
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gcc	ccaag	gag g	gccag	gaact	g to	atac	cacto	gct	caco	gag	agac	tcta	acg t	ggga	agggcc	120
cate	gataa	aac a	agcaa	aaggg	gc aa	tcct	gegg	g tta	acago	gcgt	tgcc	gcgc	aa	gcggt	gtttt	180
caco	cacca	agc a	atggg	ggaat	a co	atga	acgto	g tta	acato	aaa	gccc	ttgc	ag d	gtgt	aaagc	240
cgca	aggga	atc 9	gtgga	accc	g to	atgo	tggt	gtg	gtgga	gac	gaco	tggt	.cg t	cato	tcgga	300
gago	ccag	ggt a	aacga	aggag	ggįad	gago	gaaa	a cct	gaga	gct	,					340
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Ser 1	Thr	Val	Thr	Glu 5	Arg	Asp	Ile	Arg	Thr 10	Glu	Glu	Ser	Ile	Tyr 15	Gln	
Ala	Cys	Ser	Leu 20	Pro	Gln	Glu	Ala	Arg 25	Thr	Val	Ile	His	Ser 30	Leu	Thr	
Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	Ile	Asn	Ser	Lys 45	Gly	Gln	Ser	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Phe 60	Thr	Thr	Ser	Met	

Gly 65	Asn	Thr	Met	Thr	Cys 70	Tyr	Ile	Lys	Ala	Leu 75	Ala	Ala	Cys	Lys	Ala 80	
Ala	Gly	Ile	Val	Asp 85	Pro	Val	Met	Leu	Val 90	Сув	Gly	Asp	Asp	Leu 95	Val	
Val	Ile	Ser	Glu 100	Ser	Gln	Gly	Asn	Glu 105	Glu	Asp	Glu	Arg	Asn 110	Leu	Arg	
Ala																
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tgaa	ccgg	gag g	gccag	ggaaa	ag to	gatct	ccto	c cct	tcacc	gag	cggo	ttta	ict g	gcgga	aggccc	120
tato	gttta	aac a	agcaa	gggg	g co	cagt	gtgg	g tta	atcgo	cgt	tgc	gtgo	ca g	gtgga	gttct	180
gcct	acca	agc 1	ttgg	gcaac	ca ca	atca	actt	g tta	acato	aag	gçca	caac	gg (	ccgcg	gaaggc	240
cgca	ggc	ctc o	eggaa	cccc	gg ac	tttc	ttgt	ctg	gegga	gat	gato	tggt	.cg t	ggtg	gctga	300
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<400	)> 2	218														
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Cys	Cys	Asn	Leu 20	Glu	Pro	Glu	Ala	Arg 25	Lys	Val	Ile	Ser	30	Leu	Thr	
Glu	Arg	Leu 35	Tyr	Cys	Gly	Gly	Pro 40	Met	Pḥe	Asn	Ser	Lys 45	Gly	Ala	Gln	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Leu 60	Pro	Thr	Ser	Phe	

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Gly Asn Thr Ile Thr Cys Tyr Ile Lys Ala Thr Thr Ala Ala Lys Ala
                   70
                                        75
65
Ala Gly Leu Arg Asn Pro Asp Phe Leu Val Cys Gly Asp Asp Leu Val
Val Val Ala Glu Ser Asp Gly Val Asp Glu Asp Arg Ala Ala Leu Arg
           100
                                105
Ala
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Arg Ser Glu Gly Arg Thr Ser Trp Ala Gln
<210> 220
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Arg Ser Glu Gly Arg Thr Ser Trp Ala Gln
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Arg Thr Glu Gly Arg Thr Ser Trp Ala Gln
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<211> 629
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<400> 222

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<210> 223

<211> 209

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<213> hepatitis C virus

<400> 223

Asp Phe Trp Glu Ser Val Phe Thr Gly Leu Thr His Ile Asp Ala His 1 5 10 15

Phe Leu Ser Gln Thr Lys Gln Gln Gly Leu Asn Phe Ser Phe Leu Thr 20 25 30

Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro Pro Ser 35 40 45

Trp Asp Glu Met Trp Lys Cys Leu Val Arg Leu Lys Pro Thr Leu His 50 55 60

Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln Asn Glu Ile 65 70 75 80

Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser Ala 85 90 95

Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu Gly Gly Val Leu 100 105 110

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Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys Val Val Ile Val
                            120
        115
Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val Pro Asp Lys Glu
    130
                       135
Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala
                   150
                                        155
145
Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Glu Lys
                                    170
Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln Gln Ala Val Ile Glu
           180
Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala Phe Trp His Lys
                            200
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His
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Ile His Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile
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<211> 12
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Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile
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<211> 12 <212> PRT

<400> 226

<213> hepatitis C virus.

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Val Asn Tyr Arg Asn Ala Ser Gly Val Tyr His Ile
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<400> 227
Val Asn Tyr His Asn Thr Ser Gly Ile Tyr His Leu
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Gln His Tyr Arg Asn Ala Ser Gly Ile Tyr His Val
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<211> 12
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Gln His Tyr Arg Asn Val Ser Gly Ile Tyr His Val
<210> 230
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Ile His Tyr Arg Asn Ala Ser Asp Gly Tyr Tyr Ile
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Leu Gln Val Lys Asn Thr Ser Ser Ser Tyr Met Val

<400> 232

<400> 233

5 <210> 232 <211> 11 <212> PRT <213> hepatitis C virus

Val Trp Gln Leu Arg Ala Ile Val Leu His Val . 1

<210> 233 <211> 11 <212> PRT <213> hepatitis C virus

Val Tyr Glu Ala Asp Tyr His Ile Leu His Leu

<210> 234 <211> 11 <212> PRT <213> hepatitis C virus <400> 234

Val Tyr Glu Thr Asp Asn His Ile Leu His Leu 5

<210> 235 <211> 11 <212> PRT <213> hepatitis C virus <400> 235

Val Tyr Glu Thr Glu Asn His Ile Leu His Leu

<210> 236 <211> 11 <212> PRT <213> hepatitis C virus <400> 236

Val Phe Glu Thr Val His His Ile Leu His Leu

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Val Phe Glu Thr Glu His His Ile Leu His Leu
 1 5
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 <213> hepatitis C virus
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Val Phe Glu Thr Asp His His Ile Met His Leu
 1 5
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<211> 11
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 <213> hepatitis C virus
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 Val Tyr Glu Thr Glu Asn His Ile Leu His Leu
 1 5
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 Val Tyr Glu Ala Asp Ala Leu Ile Leu His Ala
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 Val Lys Thr Gly Asn Gln Ser Arg Cys Trp Val Ala Leu
 1 5 10
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 Val Arg Thr Gly Asn Gln Ser Arg Cys Trp Val Ala Leu
 1 5 10
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 Val Lys Thr Gly Asn Gln Ser Arg Cys Trp Ile Ala Leu
               5
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 Val Lys Thr Gly Asn Gln Ser Arg Cys Trp Ile Ala Leu
               5
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  <211> 13
  <212> PRT
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Val Lys Thr Gly Asn Ser Val Arg Cys Trp Ile Pro Leu
       5
<210> 248
<211> 13
<212> PRT
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Val Lys Thr Gly Asn Val Ser Arg Cys Trp Ile Ser Leu
1 5
<210> 249
<211> 13
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Val Arg Lys Asp Asn Val Ser Arg Cys Trp Val Gln Ile
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<210> 250
<211> 10
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Ala Pro Ser Phe Gly Ala Val Thr Ala Pro
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Val Ser Gln Pro Gly Ala Leu Thr Lys Gly
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<212> PRT
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 Val Lys Tyr Val Gly Ala Thr Thr Ala Ser
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 Ala Gln His Leu Asn Ala Pro Leu Glu Ser
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 Ser Pro Tyr Val Gly Ala Pro Leu Glu Pro
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 Ser Pro Tyr Ala Gly Ala Pro Leu Glu Pro
 1 5
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<400> 257

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Ala Pro Tyr Leu Gly Ala Pro Leu Glu Ser
             5
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Ala Pro Tyr Val Gly Ala Pro Leu Glu Ser
              5
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Asn Val Pro Tyr Leu Gly Ala Pro Leu Thr Ser
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Ala Pro His Leu Arg Ala Pro Leu Ser Ser
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Ala Pro Tyr Leu Gly Ala Pro Leu Thr Ser
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Ser Pro Gln His His Lys Phe Val Gln Asp
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<213> hepatitis C virus
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Arg Pro Arg Arg Leu Trp Thr Thr Gln Glu
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<211> 10
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<400> 266
Pro Pro Arg Ile His Glu Thr Thr Gln Asp
                5
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<211> 14
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Thr Ile Ser Tyr Ala Asn Gly Ser Gly Pro Ser Asp Asp Lys
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<210> 268

hepatitis C virus

268 <400>

Ser Arg Arg Gln Pro Ile Pro Arg Ala Arg Arg Thr Glu Gly Arg Ser

Trp Ala Gln

269 <210>

<211> 1443

<212> DNA

hepatitis C virus

<400> 269

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<211> 481

<212> PRT

<213> hepatitis C virus

<400> 270

Thr Ile Thr Thr Gly Ala Ser Ile Thr Tyr Ser Thr Tyr Gly Lys Phe 1 5 10 15

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Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly Ile Gly Thr 35 40 45

Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Ala 50 55 60

Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu 65 70 75 80

Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr Gly Arg Ala 85 90 95

Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile Phe Cys His
100 105 110

Ser Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr Ser Leu Gly 115 120 125

Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala Val Ile Pro 130 135 140 Thr Ala Gly Asp Val Val Cys Ser Thr Asp Ala Leu Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile Glu Thr Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Arg Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala Gly Glu Arg Pro Ser Asp Met Phe Asp Ser Val Val Leu Cys Glu Cys Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr Val Arg Leu Arg Ala Tyr Ile Asn Thr Pro Gly Leu Pro Val Cys Gln Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Glu Asn Phe Pro Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro Pro Pro Ser Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro Thr Leu Thr 

Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln Asn Glu Ile

Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser Ala

Asp Leu Glu Val Ile Thr Ser Thr Trp Val Leu Val Gly Gly Val Val 370 380

Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val Ala Ile Val 385 390 395 400

Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu 405 410 415

Ala Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser Ala Ser Leu 420 425 430

Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe Lys Glu Lys
435
440
445

Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu Thr Leu Lys 450 455 460

Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe Trp Ala Thr 465 470 475 480

Tyr

<210> 271

<211> 5

<212> PRT

<213> hepatitis C virus

<400> 271

Leu Glu Trp Arg Asn 5

<210> 272

<211> 20

<212> PRT

<213> hepatitis C virus

<400> 272

Leu Ser Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu Val Leu Tyr Arg 1 5 10 15

Glu Phe Asp Glu

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 phe Lys Gln Lys
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  Ser Arg Gln Ala
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    Glu Phe Asp Glu
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      Ala Ser His Leu Pro Tyr Ile Glu Gln Gly Met Gln Leu Ala Glu Gln
      <400> 276
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Phe Lys Gln Lys
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Thr Lys Gln Ala
           20
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<211> 20
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<400> 278
Val Asn Gln Arg Ala Val Val Ala Pro Asp Lys Glu Val Leu Tyr Glu
               5
Ala Phe Asp Glu
           20
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<211> 20
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Ala Ser Arg Ala Ala Leu Ile Glu Glu Gly Gln Arg Ile Ala Glu Met
Leu Lys Ser Lys
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Phe Lys Glu Lys